

Methylmercury Sorption, Uptake, and Degradation by Methanotrophs

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Microbial methylation and demethylation are two competing processes controlling the net production and bioaccumulation of neurotoxic methylmercury (CH_3Hg^+) in natural aquatic environments. While numerous studies have characterized the basis of mercury methylation, little attention has been given to the sorption, uptake, and degradation of CH_3Hg^+ by methanotrophs, despite their ubiquitous presence in the environment. We report that some methanotrophs such as *Methylosinus trichosporium* OB3b and *Methylocystis* SB2 can adsorb, take up, and degrade CH_3Hg^+ rapidly, whereas others such as *Methylococcus capsulatus* Bath and *Methylovulum miyakonense* can take up but not degrade CH_3Hg^+ . Demethylation by *M. trichosporium* OB3b increases with increasing CH_3Hg^+ concentrations but was abolished in mutants deficient in the synthesis of methanobactin, a metal-binding compound used by some methanotrophs such as *M. trichosporium* OB3b. Further, addition of methanol (> 5 mM) as a competing one-carbon (C1) substrate inhibits demethylation, suggesting that CH_3Hg^+ degradation by methanotrophs may involve an initial bonding of CH_3Hg^+ by methanobactin followed by cleavage of the C–Hg bond in CH_3Hg^+ by the methanol dehydrogenase. These observations demonstrate a previously unknown but potentially important mechanism for CH_3Hg^+ detoxification and suggest possible broader involvement of C1-metabolizing aerobes in the degradation and cycling of toxic CH_3Hg^+ in the environment.